

1/31



OICE

RAW SEQUENCE LISTING

DATE: 02/06/2002

PATENT APPLICATION: US/10/027,199

TIME: 09:52:25

Input Set : N:\Crif3\RULE60\10027199.raw

Output Set: N:\CRF3\02062002\J027199.raw

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1 <110> APPLICANT: Kwon, Byoung
2 <120> TITLE OF INVENTION: NEW RECEPTOR AND RELATED PRODUCTS AND
3   METHODS
4 <130> FILE REFERENCE: 740.013US2
5 <140> CURRENT APPLICATION NUMBER: 10/027,199
6 <141> CURRENT FILING DATE: 2001-12-20
7 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/955,572
W--> 8 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-22
9 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/461,652
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11 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/122,796
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13 <160> NUMBER OF SEQ ID NOS: 12
14 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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17 <211> LENGTH: 838
18 <212> TYPE: DNA
19 <213> ORGANISM: Homo sapiens
20 <400> SEQUENCE: 1
21   aatcagcttt gctagtatca tacctgtgcc agatttcac atgggaaaca gctgttacaa      60
22   catagtagcc actctgttgc tggctctcaa ctttgagagg acaagatcat tgcaggatcc      120
23   ttgtagtaac tgcccagctg gtacattctg tgataataac aggaatcaga ttgacagtcc      180
24   ctgtcctcca aatagtttct ccagcgcagg tggacaaagg acctgtgaca tatgcaggca      240
25   gtgtaaaggt gttttcagga ccaggaagga gtgttctcc accagcaatg cagagtgtga      300
26   ctgcactcca gggtttcaact gcctgggggc aggatgcagc atgtgtgaac aggatgttaa      360
27   acaagggtcaa gaactgacaa aaaaagggtg taaagactgt tgctttggga catttaacga      420
28   tcagaaacgt ggcactctgt gacctggac aaactgttct ttggatggaa agtctgtgct      480
29   tgtgaatggg acgaaggaga gggacgtggt ctgtggacca tctccagctg acctctctcc      540
30   gggagcctcc tctgtgaccc cgctgcccc tgcgagagag ccaggacact ctccgcagat      600
31   catctccttc tttcttgccg tgacgtcgac tgcgttgctc ttctgtgtgt tcttctctac      660
32   gctccgtttc tctgttggtt aacggggcag aaagaaactc ctgtatatat tcaaacaacc      720
33   atttatgaga ccagtacaaa ctactcaaga ggaagatggc tgtagctgcc gatttccaga      780
34   agaagaagaa ggaggatgtg aactgtgaaa tggaaagtaa tagggctgtt gggacttt      838
36 <210> SEQ ID NO: 2
37 <211> LENGTH: 255
38 <212> TYPE: PRT
39 <213> ORGANISM: Homo sapiens
40 <400> SEQUENCE: 2
41   Met Gly Asn Ser Cys Tyr Asn Ile Val Ala Thr Leu Leu Leu Val Leu
42   1             5             10             15
43   Asn Phe Glu Arg Thr Arg Ser Leu Gln Asp Pro Cys Ser Asn Cys Pro
44   20             25             30
45   Ala Gly Thr Phe Cys Asp Asn Asn Arg Asn Gln Ile Cys Ser Pro Cys

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46          35          40          45
47    Pro Pro Asn Ser Phe Ser Ser Ala Gly Gly Gln Arg Thr Cys Asp Ile
48          50          55          60
49    Cys Arg Gln Cys Lys Gly Val Phe Arg Thr Arg Lys Glu Cys Ser Ser
50    65          70          75          80
51    Thr Ser Asn Ala Glu Cys Asp Cys Thr Pro Gly Phe His Cys Leu Gly
52          85          90          95
53    Ala Gly Cys Ser Met Cys Glu Gln Asp Cys Lys Gln Gly Gln Glu Leu
54          100          105          110
55    Thr Lys Lys Gly Cys Lys Asp Cys Cys Phe Gly Thr Phe Asn Asp Gln
56          115          120          125
57    Lys Arg Gly Ile Cys Arg Pro Trp Thr Asn Cys Ser Leu Asp Gly Lys
58          130          135          140
59    Ser Val Leu Val Asn Gly Thr Lys Glu Arg Asp Val Val Cys Gly Pro
60    145          150          155          160
61    Ser Pro Ala Asp Leu Ser Pro Gly Ala Ser Ser Val Thr Pro Pro Ala
62          165          170          175
63    Pro Ala Arg Glu Pro Gly His Ser Pro Gln Ile Ile Ser Phe Phe Leu
64          180          185          190
65    Ala Leu Thr Ser Thr Ala Leu Leu Phe Leu Leu Phe Phe Leu Thr Leu
66          195          200          205
67    Arg Phe Ser Val Val Lys Arg Gly Arg Lys Lys Leu Leu Tyr Ile Phe
68          210          215          220
69    Lys Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln Glu Glu Asp Gly
70    225          230          235          240
71    Cys Ser Cys Arg Phe Pro Glu Glu Glu Glu Gly Gly Cys Glu Leu
72          245          250          255

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74 <210> SEQ ID NO: 3

75 <211> LENGTH: 20

76 <212> TYPE: DNA

77 <213> ORGANISM: Homo sapiens

78 <400> SEQUENCE: 3

79 ttytgymgaa artayaaycc

20

81 <210> SEQ ID NO: 4

82 <211> LENGTH: 20

83 <212> TYPE: DNA

84 <213> ORGANISM: Homo sapiens

85 <400> SEQUENCE: 4

86 ttytctstsc a htggtggaca

20

88 <210> SEQ ID NO: 5

89 <211> LENGTH: 20

90 <212> TYPE: DNA

91 <213> ORGANISM: Homo sapiens

92 <400> SEQUENCE: 5

93 cccargswrc aggttytrca

20

95 <210> SEQ ID NO: 6

96 <211> LENGTH: 20

97 <212> TYPE: DNA

98 <213> ORGANISM: Homo sapiens

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Input Set : N:\Crif3\RULE60\10027199.raw

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100      ttytgrtctt traatgttcc                                20
102 <210> SEQ ID NO: 7
103 <211> LENGTH: 25
104 <212> TYPE: DNA
105 <213> ORGANISM: Homo sapiens
106 <400> SEQUENCE: 7
107      aataagcttt gctagtatca tacct                                25
109 <210> SEQ ID NO: 8
110 <211> LENGTH: 30
111 <212> TYPE: DNA
112 <213> ORGANISM: Homo sapiens
113 <400> SEQUENCE: 8
114      ttaagatctc tgccgagagt gtcctggctc                                30
116 <210> SEQ ID NO: 9
117 <211> LENGTH: 2350
118 <212> TYPE: DNA
119 <213> ORGANISM: Mus musculus
120 <220> FEATURE:
121 <221> NAME/KEY: unsure
122 <222> LOCATION: (1253)...(1255)
123 <223> OTHER INFORMATION: (a or g or c or t/u)
124 <400> SEQUENCE: 9
125      atgtccatga actgctgagt ggataaacag cacgggatat ctctgtctaa aggaatatta                                60
126      ctacaccagg aaaaggacac attcgacaac aggaaaggag cctgtcacag aaaaccacag                                120
127      tgtcctgtgc atgtgacatt tcgccatggg aaacaactgt tacaacgtgg tggtcattgt                                180
128      gctgctgcta gtgggctgtg agaagggtggg agccgtgcag aactcctgtg ataactgtca                                240
129      gcctggtaact ttctgcagaa aatacaatcc agtctgcaag agctgccctc caagtacctt                                300
130      ctccagcata ggtggacagc cgaactgtaa catctgcaga gtgtgtgcag gctatttcag                                360
131      gttcaagaag ttttgcctct ctaccacaaa cgcggagtgt gagtgcattg aaggattcca                                420
132      ttgcttgggg ccacagtgca ccagatgtga aaaggactgc aggcctggcc aggagctaac                                480
133      gaagcagggt tgcaaaacct gtagcttggg aacatttaat gaccagaacg gtactggcgt                                540
134      ctgtcgaccc tggacgaact gctctctaga cggaaggctc gtgcttaaga ccgggaccac                                600
135      ggagaaggac gtggtgtgtg gacccccctg ggtgagcttc tctcccagta ccaccatttc                                660
136      tgtgactcca gagggaggac caggagggca ctcttgcag gtccttacct tgttctggc                                720
137      gctgacatcg gctttgtgtc tggccctgat ctccattact ctctgtttct ctgtgtctca                                780
138      atggatcagg aaaaaattcc cccacatatt caagcaacca tttaagaaga ccactggagc                                840
139      agtcaagag gaagatgctt gtagctgccg atgtccacag gaagaagaag gagggaggag                                900
140      aggtatgag ctgtgatgta ctatcctagg agatgtgtgg gccgaaaccg agaagcacta                                960
141      ggacccccacc atcctgtgga acagcacaag caacccccacc acctgttct tacacatcat                                1020
142      cctagatgat gtgtgggcgc gcacctcatc caagtctctt ctaacgctaa catatttgtc                                1080
143      tttacctttt ttaaattctt ttttaaatth aaattttatg tgtgtgagtg ttttgcctgc                                1140
144      ctgtatgcac acgtgtgtgt gtgtgtgtgt gtgacactcc tgatgcctga ggaggtcaga                                1200
W--> 145      agacaaaggg ttggttccat aagaactgga gttatggatg gctgtgagcc ggnnngatag                                1260
146      gtcgggacgg agacctgtct tcttatttta acgtgactgt ataataaaaa aaaaatgata                                1320
147      ttccgggaat tgtagagatt gtcctgacac ccttctagtt aatgatctaa gaggaattgt                                1380
148      tgatacgtag tatactgtat atgtgtatgt atatgtatat gtatatataa gactctttta                                1440
149      ctgtcaaagt caacctagag tgtctgggta ccaggtcaat tttattggac attttacgtc                                1500
150      acacacacac acacacacac acacacacgt ttatactacg tactgtttac ggtattctac                                1560

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151 gtcataataat gggatagggg aaaaggaac caaagagtga gtgatattat tggaggagt 1620
152 acagactacc ccttctgggt acgtagggac agacctcctt cggactgtct aaaactcccc 1680
153 ttagaagtct cgtcaagttc cggacgaag aggacagagg agacacagtc cgaagaagtta 1740
154 ttttccggc aaatccttcc cctgttctgt gacactccac ccttgtgga caattgagt 1800
155 tcatccttgc gccggaagggt caggtggtac ccgtctgtag gggcggggag acagagccgc 1860
156 gggggagcta cgagaatcga ctacagggc gcccgggct togc aaatga aactttttta 1920
157 atctcacaag ttctgtccgg gtctggcgga cctatggcgt cgtaccttat taacttatcc 1980
158 tggcgccaag ataaaaaac caaaagcctt gactccggta ctaattctcc ctgcccgc 2040
159 ccgtaagcat aacggcgga tctccacttt aagaacctgg ccggttctg cctggtctcg 2100
160 ctttctgaaa cgttctttac aaaagtaatt agttcttctt ttcagctcc aagcttctgc 2160
161 tagtctatgg cagcatcaag gctggtatct gctacggctg accgctacgc cgcgcgaata 2220
162 agggtaactg gcggcccggt gaaggccctt tggtttcaga aacccaaggc cccctcata 2280
163 ccaacgtttc gactttgatt ctgcccggta cgtggtggtg ggtgccttag ctcttctcg 2340
164 atagttagac 2350
166 <210> SEQ ID NO: 10
167 <211> LENGTH: 256
168 <212> TYPE: PRT
169 <213> ORGANISM: Mus musculus
170 <400> SEQUENCE: 10
171 Met Gly Asn Asn Cys Tyr Asn Val Val Val Ile Val Leu Leu Leu Val
172 1 5 10 15
173 Gly Cys Glu Lys Val Gly Ala Val Gln Asn Ser Cys Asp Asn Cys Gln
174 20 25 30
175 Pro Gly Thr Phe Cys Arg Lys Tyr Asn Pro Val Cys Lys Ser Cys Pro
176 35 40 45
177 Pro Ser Thr Phe Ser Ser Ile Gly Gly Gln Pro Asn Cys Asn Ile Cys
178 50 55 60
179 Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser Thr
180 65 70 75 80
181 His Asn Ala Glu Cys Glu Cys Ile Glu Gly Phe His Cys Leu Gly Pro
182 85 90 95
183 Gln Cys Thr Arg Cys Glu Lys Asp Cys Arg Pro Gly Gln Glu Leu Thr
184 100 105 110
185 Lys Gln Gly Cys Lys Thr Cys Ser Leu Gly Thr Phe Asn Asp Gln Asn
186 115 120 125
187 Gly Thr Gly Val Cys Arg Pro Trp Thr Asn Cys Ser Leu Asp Gly Arg
188 130 135 140
189 Ser Val Leu Lys Thr Gly Thr Thr Glu Lys Asp Val Val Cys Gly Pro
190 145 150 155 160
191 Pro Val Val Ser Phe Ser Pro Ser Thr Thr Ile Ser Val Thr Pro Glu
192 165 170 175
193 Gly Gly Pro Gly Gly His Ser Leu Gln Val Leu Thr Leu Phe Leu Ala
194 180 185 190
195 Leu Thr Ser Ala Leu Leu Leu Ala Leu Ile Phe Ile Thr Leu Leu Phe
196 195 200 205
197 Ser Val Leu Lys Trp Ile Arg Lys Lys Phe Pro His Ile Phe Lys Gln
198 210 215 220
199 Pro Phe Lys Lys Thr Thr Gly Ala Ala Gln Glu Glu Asp Ala Cys Ser
200 225 230 235 240

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DATE: 02/06/2002

TIME: 09:52:25

Input Set : N:\Crf3\RULE60\10027199.raw

Output Set: N:\CRF3\02062002\J027199.raw

```

201      Cys Arg Cys Pro Gln Glu Glu Glu Gly Gly Gly Gly Tyr Glu Leu
202                                245                250                255
204 <210> SEQ ID NO: 11
205 <211> LENGTH: 24
206 <212> TYPE: PRT
207 <213> ORGANISM: Homo sapiens
208 <220> FEATURE:
209 <221> NAME/KEY: ZN_FING
210 <222> LOCATION: 2...3, 5...13, 15...17, 19...21, 23
211 <223> OTHER INFORMATION: Putative zinc finger structure
212 <400> SEQUENCE: 11
W--> 213      Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
214              1              5              10              15
W--> 215      Xaa His Xaa Xaa Xaa Cys Xaa Cys
216              20
218 <210> SEQ ID NO: 12
219 <211> LENGTH: 12
220 <212> TYPE: PRT
221 <213> ORGANISM: Homo sapiens
222 <400> SEQUENCE: 12
223      Leu Gln Asp Pro Cys Ser Asn Cys Pro Ala Gly Thr
224              1              5              10

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VERIFICATION SUMMARY

DATE: 02/06/2002

PATENT APPLICATION: US/10/027,199

TIME: 09:02:26

Input Set : N:\Crf3\RULE60\10027199.raw

Output Set: N:\CRF3\02062002\J027199.raw

L:8 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:10 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:12 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:213 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11